Supplementary Table 1. Summary of Subject Genotyping Information

Analysis Set/Derivation	subjects	Platform	Quality ^a	IBD^b	Gender
Discovery EOMG Sweden	189	Illumina 370K	10	1	2
Discovery EOMG England	121	Illumina 370K	3	1	0
Discovery EOMG France	118	Illumina 370K	0	3	0
Discovery EOMG Norway	26	Illumina 370K	0	0	0
Total Discovery EOMG Cases	454				
Replication EOMG Swedish	10	Illumina 610K	0	0	0
Replication EOMG English	128	Illumina 610K	10	0	5
Replication EOMG German	26	Illumina 610K	0	0	1
Replication EOMG Netherlands	46	Illumina 610K	0	0	0
Replication EOMG Norway	76	Illumina 610K	1	4	5
Total Replication EOMG Cases	286				
Discovery Controls	subjects	Platform	Quality ^a	IBD ^b	Gender
Discovery Controls Sweden	subjects 570	Platform Illumina 550K	Quality^a 0	IBD ^b 0	Gender 0
			-		
Sweden	570	Illumina 550K	0	0	0
Sweden British 1958 Blood Bank	570 1355	Illumina 550K Illumina 550K	0 19	0 13	0 0
Sweden British 1958 Blood Bank New York Cancer Project	570 1355 1834	Illumina 550K Illumina 550K Illumina 550K	0 19 10	0 13 10	0 0 4
Sweden British 1958 Blood Bank New York Cancer Project Norway ^f	570 1355 1834 123	Illumina 550K Illumina 550K Illumina 550K Illumina 550K	0 19 10 4	0 13 10 0	0 0 4 0
Sweden British 1958 Blood Bank New York Cancer Project Norway ^f Childrens Hospital of Philadelphia	570 1355 1834 123 1453	Illumina 550K Illumina 550K Illumina 550K Illumina 550K Illumina 550K	0 19 10 4 23	0 13 10 0 47	0 0 4 0
Sweden British 1958 Blood Bank New York Cancer Project Norway ^f Childrens Hospital of Philadelphia NIH Laboratory of Neurogenetics	570 1355 1834 123 1453 170	Illumina 550K Illumina 550K Illumina 550K Illumina 550K Illumina 550K	0 19 10 4 23	0 13 10 0 47	0 0 4 0
Sweden British 1958 Blood Bank New York Cancer Project Norway ^f Childrens Hospital of Philadelphia NIH Laboratory of Neurogenetics Total	570 1355 1834 123 1453 170 5505	Illumina 550K Illumina 550K Illumina 550K Illumina 550K Illumina 550K Illumina 550K	0 19 10 4 23 0	0 13 10 0 47 0	0 0 4 0 0

a. Less than 95% complete genotypes

b. Exclusion for relatedness based on identity by state analyses (PI^>0.15 assessed using PLINK IBD

c. >10% nonEuropean admixture (see Supplementary Methods)

d. Based on PCA analyses (see Supplementary Methods)

e. Final set after exclusion or after best 4:1 match for controls (see Supplementary Methods).

f. Samples collected in Norway and genotyped in current study at North Shore.

g. Samples collected and genotyped in Sweden for population controls in myocardial infarction study.

non-European ^c	4 SD PCA ^d	Final Set ⁶
9	5	162
3	1	113
16	0	99
0	0	26
		400
0	0	10
2	4	107
0	0	25
2	1	43
0	2	64
		249

non-European^c 4 SD PCA^d 4:1 match cases^e

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5	5	535
18	15	292
115	20	310
0	4	98
169	10	305
16	5	60
		1600
6	23	309
38	13	687
		996

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